

<u>DOMAINS</u>: I - Cytoplasmic Domain; II - Transmembrane Domain; III - Proximal Extracellular Domain; IV - Distal Extracellular Domain (putative soluble form)

Figure 1

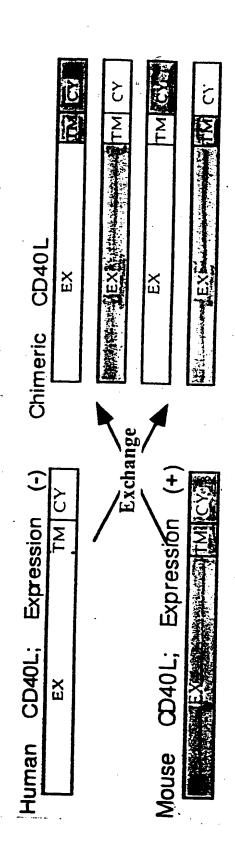


Figure 2

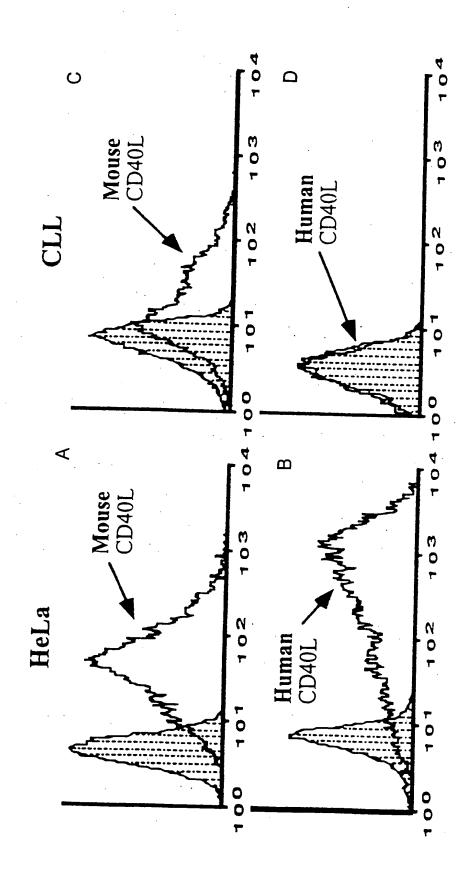


Figure 3

### **NONINFECTED**

### +mCD40-L ADENOVIRUS

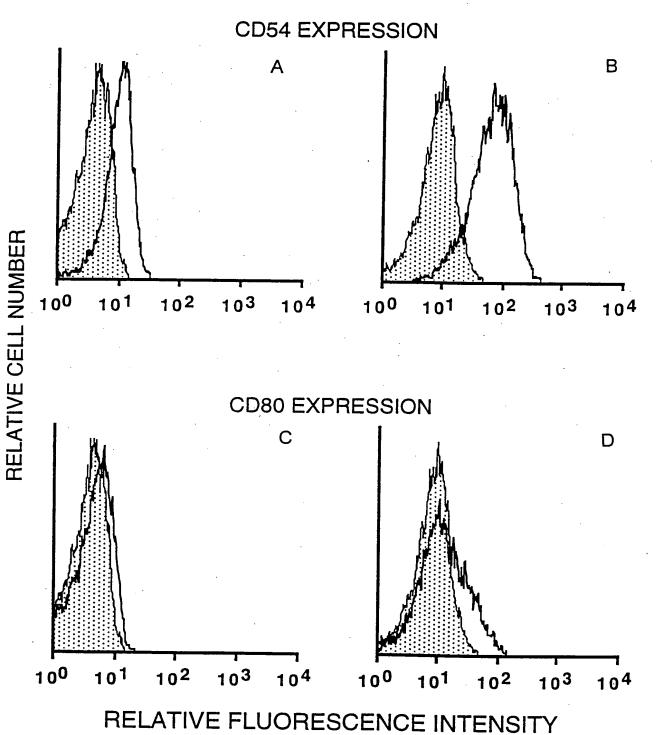


Figure 4

-

# Allogenic T cell response to CLL cells transfected with adeno-mCD40L

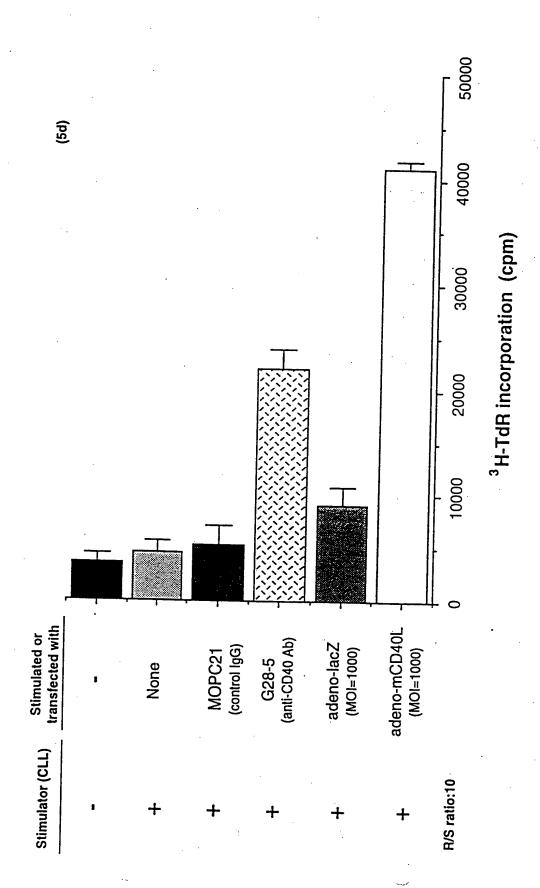


Figure 5

## Production of IFNy by allogenic T lymphocytes stimulated with CLL B cells

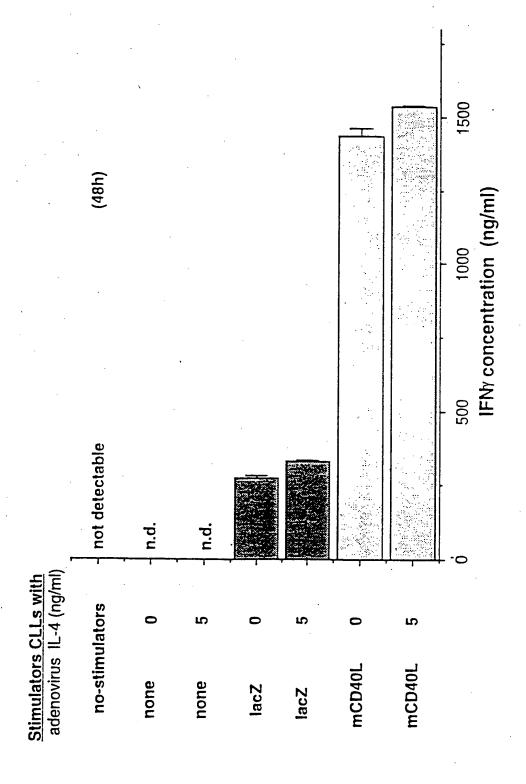


Figure 6

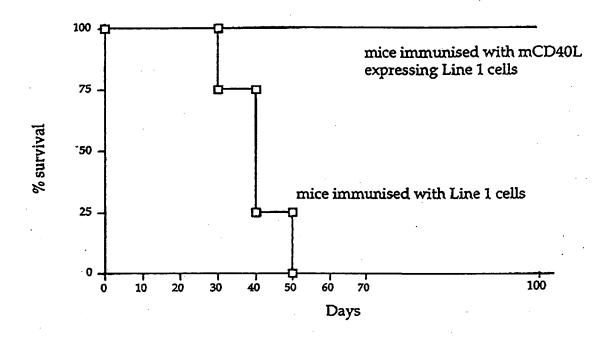


Figure 7

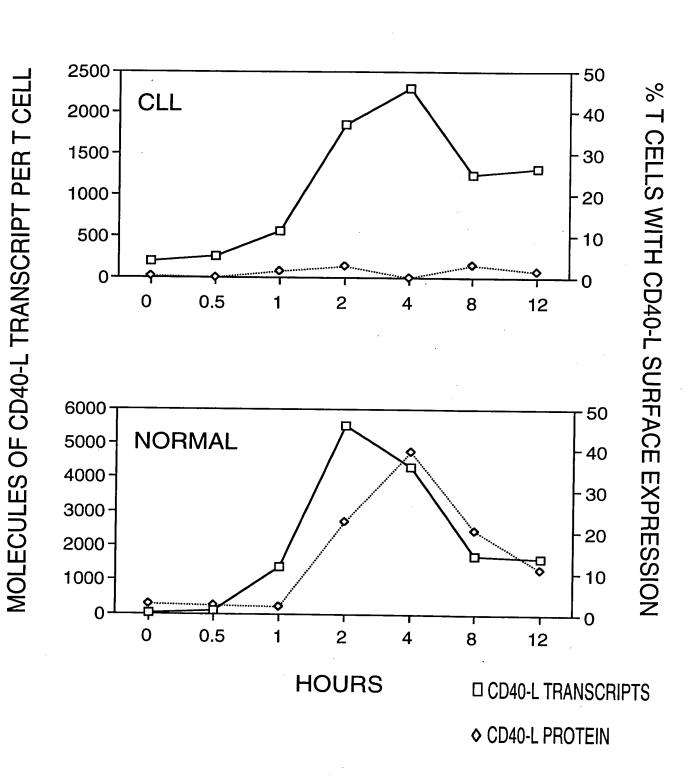


Figure 8

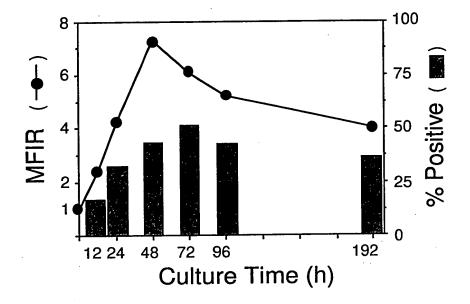


Figure 9

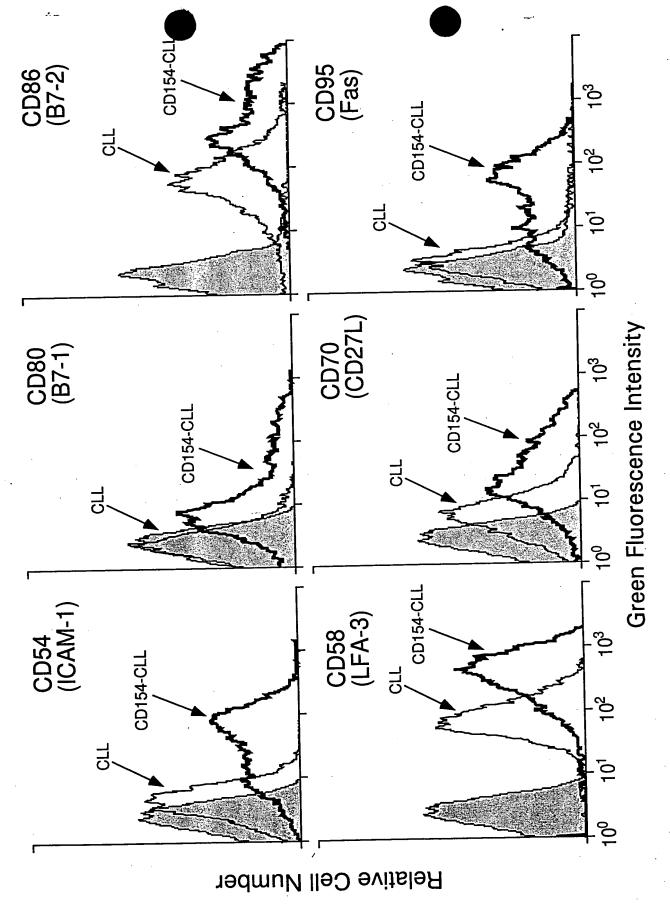


Figure 10

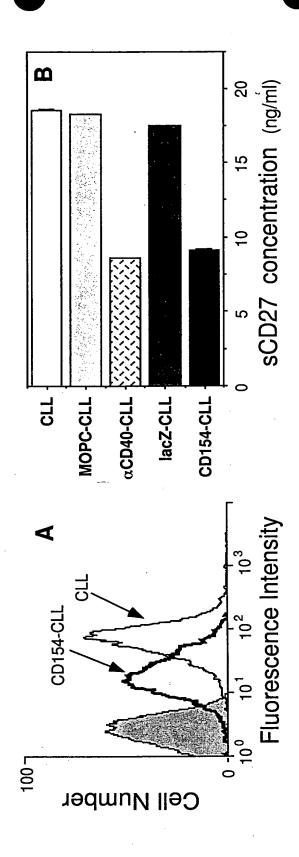


Figure 11

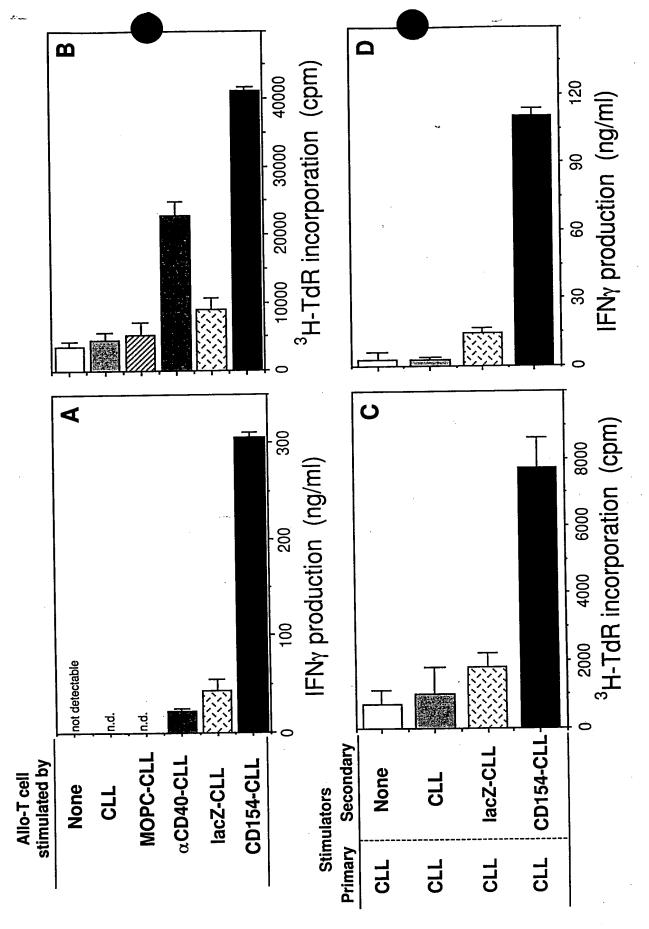


Figure 12

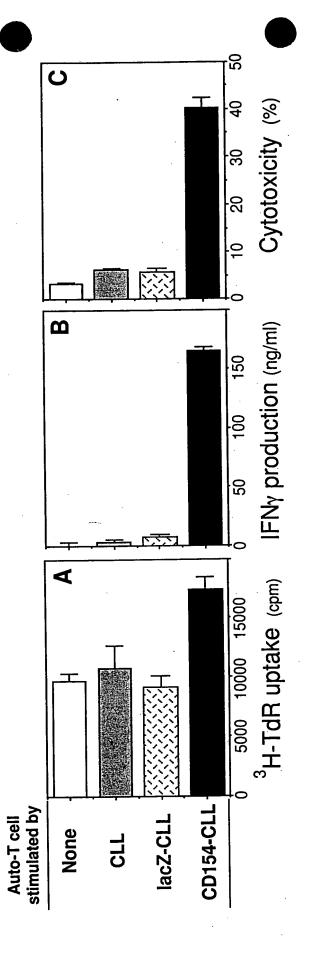


Figure 13

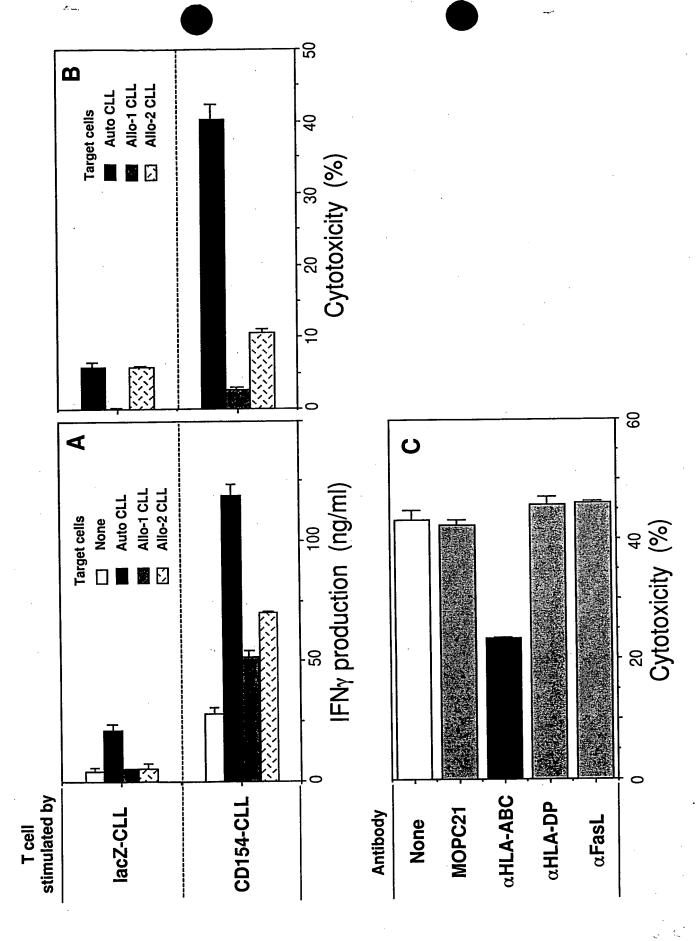


Figure 14

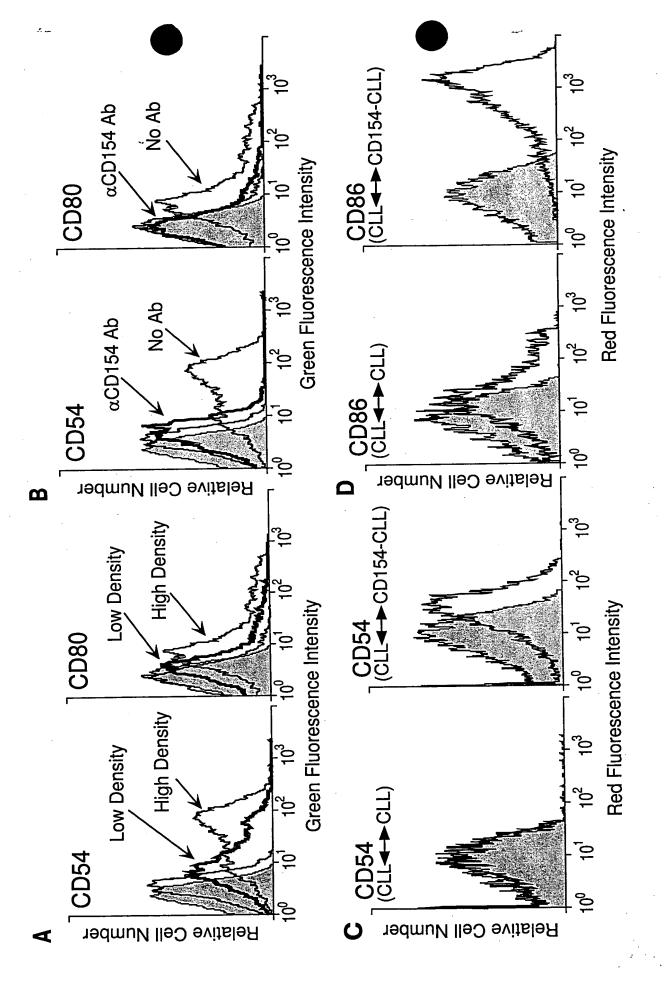


Figure 15

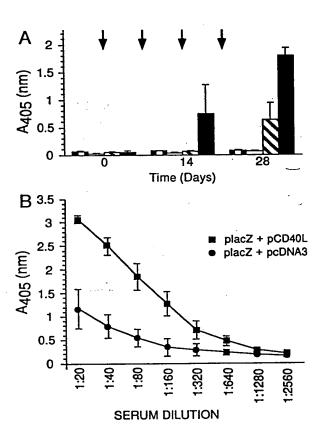
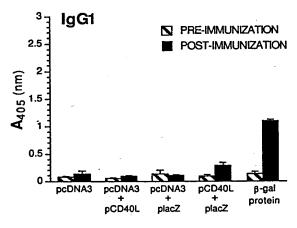


Figure 16



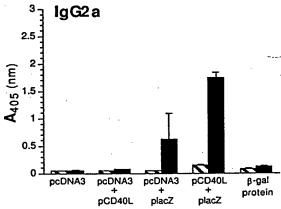


Figure 17

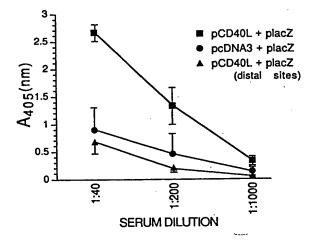


Figure 18

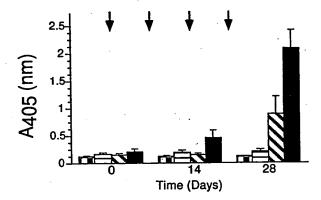


Figure 19

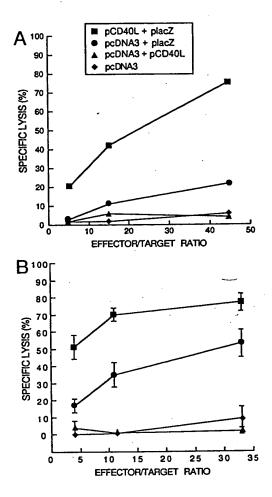
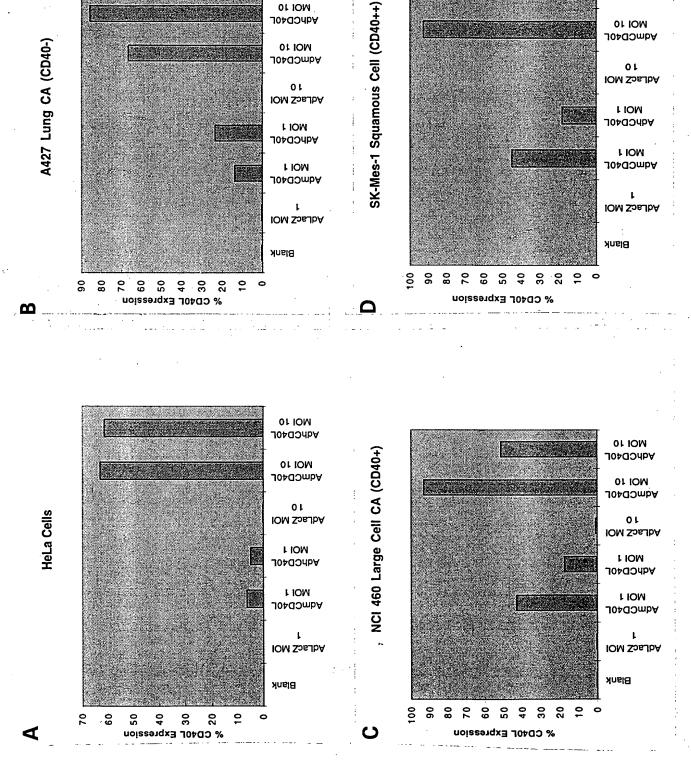
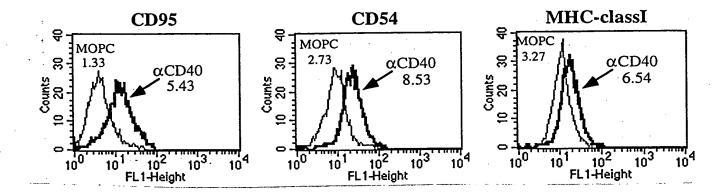


Figure 20



igure 21

WOI 10 VQPCD40F



В

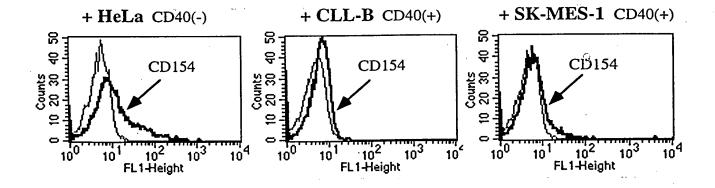


Figure 22

Figure 23

## Gene Therapy of Leukemia

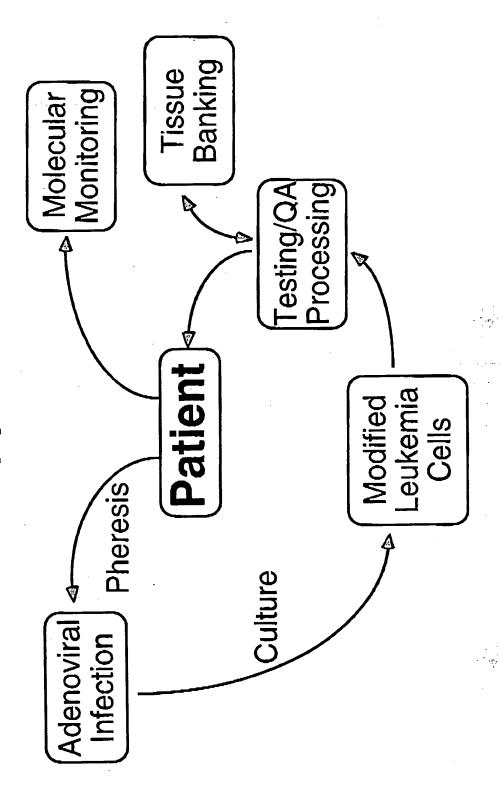


Figure 24

	MOOPFNYPYPOIYWVDSSASSPWAPPGTVLPCPTSVPRRPGORRPPPPPPP	
1	THE THEORY OF THE PROPERTY OF	50
51	PPPLPPPPPPPPPLPLPLPPLKKRGNHSTGLCLLVMFFMVLVALVGLGLG	100
	THE PROPERTY OF THE PROPERTY O	
51	MFOLFHLOKELAELRESTSOMHTASSLEKOIGHPSPPPEKKELRKVAHLT	
101		
101	TO THE PROPERTY OF THE PROPERT	150
151	GKSNSRSMPLEWEDTYGIVLLSGVKYKKGGLVINETGLYFVYSKVYFRGQ	200
1 5 1	GKSNSRSMPLEWEDTYGIVLLSGVKYKKGGLVINETGLYFVYSKVYFRGQ	
	THE	
201		
201	SCNNLPLSHKVYMRNSKYPODLVMMEGKMMSYCTTGQMWARSSYLGAVFN	250
251	LTSADHLYVNVSELSLVNFEESOTFFGLYKL 281	
251		

	granding and a second restriction for the first of the fi	
	MOOPFNYPYPOIYWVDSSASSFWAPPGTVLPCPTSVPRRPGORRPPPPPPPPPPPPPPPPPPPPPPPPPPPPPPP	
1	MQQPFNYPYPQIYWVDSSASSPWAPPGTVLPCPTSVPRRPGQRRPPPPPP	50
51	PPPLPPPPPPPPPLPPLPPLKKRGNHSTGLCLLVMFFMVLVALVGLGLG	100
51		100
	MFOLFHLOKELAELRESTSOMHTASSLEKQIGHPSPPPEKKELRKVAHLT	
		130
151	GKSNSRSMPLEWEDTYGIVLLSGVKYKKGGLVINETGLYFVYSKVYFRGQ	200
131	GKSNSRSMPLEWEDTYGIVLLSGVKYKKGGLVINETGLYFVYSKVYFRGQ	180
201	SCNINLPLSHKVYMRNSKYPODLVMMEGKMMSYCTTGOMWARSSYLGAVFN	250
	SCHOOL DESIRED TO THE SCHOOL OF THE SCHOOL O	
	LTSADHLYVNVSELSLVNFEESOTFFGLYKL 281	
	4   4   5   4   4   5   5   5   5   5	
231	LTSADHLYVNVSELSLVNFEESQTFFGLYKL 261	

	MQQPFNYPYPQIYWVDSSASSPWAPPGTVLPCPTSVPRRPGORRPPPPPPP	50
1	MOOPFNYPYPOIYWVDSSASSPWAPPGTUPPCHILIPHILIPHILIPHILIPHILIPHILIPHILIPHI	
1		50
_	PPPLPPPPPPPPPPLPLPPLKKRGNHSTGLCLLVMFFMVLVALVGLGLG	100
<b>E</b> 1		100
101	MFQLFHLOKELAELRESTSOMHTASSLEKQQIGHPSPPPEKKELRKVAHLT	
101	MFQLF. MPEEGSGCSVRRRPYGCVLRIGHPSPPPEKKELRKVAHLT	145
151	GKSNSRSMPLEWEDTYGIVLLSGVKYKKGGLVINETGLYFVYSKVYFRGO	
146	GKSNSRSMPLEWEDTYGIVLLSGVKYKKGGLVINETGLYFVYSKVYFRGQ	
	SCNINLPLSHKVYMRNSKYPODLVMMEGKMMSYCTTGOMWARSSYLGAVFN	250
201	SCNNLPLSHKVYMRNSKYPODLOUMERKAMISTULLI IIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII	
		245
196	SCNNLPLSHKVYMRNSKYPQDDVFILLEGRATIO1010104	
	LTSADHLYVNVSELSLVNFEESOTFFGLYKL 281	
251	ITSADHLYVNVSEDSLVNFEDSOXII	
246	UTSADHLYVNVSELSUVMFEESQ11 CODIA	

### Matrix Metalloproteinase Cleavage Sites

		Cleavage ↓					
₽,	ሚ	<b>6</b>	σ.	፫	د	ŗ.	<u>o</u>
Collagenases	ases						
MMP-1	Interstitial Collager	Collagenase					
Φ.	₾.	<b>D</b> 2	ሚ	ŗ	<b>ب</b> '	Ţ	<u>ଫ</u>
Ala	Pro	Leu	Gly	Met	Arg	Met/Ala	Arg
Glv/Leu	Leu	Met/Tyr	His	ren	ren	Gly	Lys
Met Met	Ala	Val/Gly	Glu	<u>e</u>	Phe	Val	Gh
Glu	Asp	<u>e</u>	Tyr	GIn	T.	Ser	<u>е</u>
Pro	Ser	Gln/Arg	Ala	Pro	Olu	Glu	Gly
ĭ×	Glu	Asp	Phe	Phe	Ala	Phe	Ser
` <u>e</u>	ζĮ	Ölü	Gľu	Ala	Val/Gly	Arg	Glu
Thr	Arg	Ala	Asn	Tyr/Val	Ser	Pro	Ala
Arg	)			[not K,E,W]	Asn		
MMP-8	Neutrophil	Neutrophil Collagenase					
<b>₽</b>	ሚ	ሚ	ሚ	<b>Ţ</b>	<b>ب</b>	Ţ	<u>,</u>
Ala	Pro	Leu	Glu	Туr	Ala	Gly	Arg
Gly/Leu	ren	GIn	Gly/His	<u>e</u>	Leu	Met	Gin
Met	Met		Ala	ren	Trp	Ala	
Glu	•			Val			
Pro				Phe			
Tyr/lle/Thr/	/Arg		Ala				
(otherwise	same as MMI	P-1)					

Stromelys	SIIIS					
MMP-3 SI	tromelysin 1					,
<b>P</b>	_ლ	<b>~</b>	ሚ	ŗ	2,	ŗ.
Asp	Pro	Phe	Glu	Leu	Arg	Ala
Gly Ala	Ala	Leu/Met	Ala	Phe	Leu/Phe	Arg/Met
Gln/Arg						į
ren	Val	Tyr	Gln/Phe	Trp/Tyr	т Б	S S
<u>le</u>	Leu	Pro/Gly/Glu	Asn	<u>e</u>	Val	Val/lle
Glu/Val						;
Leu	Thr	<u>e</u>	His	Val	Gh	Ser/Asn
Lys	Phe	Ala	Gly	Met	His/Met	Glu/Thr
Gly/Asp						
Arg	Arg	Ser	Leu/Pro	Oln Oln	Glu/Ser/Thr	Leu
Ser/Lys/Phe						
Pro/Met	Ser/Gly		Lys/Tyr/Arg			

Pro

**₽**.

Figure 28B

Ala/Phe/Gln

<b>.</b> .⊒ .⊒	Ala			₽_	등		•	
<b>P</b> ,	Glu			Ţ	Ala	Val/Arg/Met	Gly	
<b>.</b> G. 2	Val			Ţ,	Arg	Met	Gln	
<b>g</b> =	Геп			ŗ.	<b>Leu</b>	<u>e</u>	Met	
<b>-</b> H ∃is	Leu		·	ሚ	Olu Glu	<b>Met/Ala</b>	Pro/Gln	Gly
<b>o</b> 2 <u>⊕</u>				ሚ	Leu	Gln	Val	
Stromelysin 2 P <sub>3</sub> Ala	Pro	,	Matrilysin	ሚ	Pro	Leu		
MMP-10 <b>P</b> ₄ Ara	Gly	Others	MMP-7	σ <u>,</u>	<u>e</u>	Gly	Pro	